



X-TRA-NET ChIP-seq workshop June 24-25, 2009

Strasbourg – France

**The Institute of Genetics and Molecular and Cellular Biology (IGBMC)
ILLKIRCH CEDEX**

Wednesday, June 24

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|-------------|---|
| 8.00 - 9:00 | Registration - coffee |
| 9.00 | Welcome – Introduction (Hinrich Gronemeyer) |
| 9.10 | Illumina – Technology and Applications |
| 10.00 | Illumina – Pipeline and data analysis |
| 10.45 | Coffee break |
| 11.00 | Bernard Jost - The IGBMC core facility – Technology and Applications
<i>(Sample prep and sample flow in the core facility)</i> |
| 11.45 | Stephanie Le Gras - The IGBMC core facility: Primary data treatment
and filtering (IT requirements, data handling and storage) |
| 12.30 | CLCBio – Software introduction |
| 13.00 | Lunch |
| 14.00 | ChIP-seq Workshop – Introduction (Hinrich Gronemeyer)

Susanne Mandrup – Introduction to hands-on training data set |
| 14.30 | CLCBio – Hands-on |
| 16.30 | 4-6 Short talks (15 min each) selected from participant abstracts |
| 18.00 | End of day 1 |
| 19.00 | Dinner |

Thursday, June 25

9.00	Welcome – Day 2 (Hinrich Gronemeyer)
9.00	Danilo Ceschin (IGBMC) - Bioinformatic analysis of ChIP-seq data
9.45	Blaise Alako (Radboud University) - Publicly available software
10.30	Coffee break
11.00	Albin Sandelin (BRIC, University of Copenhagen) - Regulatory meta-analysis of ChIP and tag data
11.45	Fiona Nielsen (Radboud University) – CATCH
12.15	Genomatix – Software introduction
12.45	Lunch
13.45	Hands-on practice Day 2 Genomatix – Hands-on
16.00	Public available software – Software introduction
16.30	Public available software – Hands-on
18.30	End of day 2
20.00	Dinner/Departure